

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/590,552
Source: IFWP
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IFWP

RAW SEQUENCE LISTING

DATE: 09/01/2006

PATENT APPLICATION: US/10/590,552

TIME: 12:08:10

Input Set : F:\SEQUENCE LISTING.txt

Output Set: N:\CRF4\09012006\J590552.raw

3 <110> APPLICANT: ITO, Yoshitaka
 4 TAKAMIZAWA, Kazuhiro
 5 IWAHASHI, Hitoshi
 7 <120> TITLE OF INVENTION: METHOD OF JUDGING BIOLOGICAL ACTIVITY IN BIOREMEDIATION SITE

AND

8 POLYNUCLEOTIDE FOR DETECTING MICROORGANISM TO BE USED THEREIN

10 <130> FILE REFERENCE: 10873.1940USWO

C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/590,552

13 <141> CURRENT FILING DATE: 2006-08-23

15 <150> PRIOR APPLICATION NUMBER: PCT/JP2005/003175

16 <151> PRIOR FILING DATE: 2005-02-25

19 <150> PRIOR APPLICATION NUMBER: JP application No.2004-50082

20 <151> PRIOR FILING DATE: 2004-02-25

22 <150> PRIOR APPLICATION NUMBER: JP application No.2004-50083

23 <151> PRIOR FILING DATE: 2004-02-25

25 <160> NUMBER OF SEQ ID NOS: 118

27 <170> SOFTWARE: PatentIn version 3.3

29 <210> SEQ ID NO: 1

30 <211> LENGTH: 742

31 <212> TYPE: DNA

32 <213> ORGANISM: Dehalospirillum multivorans

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43	gcgcctgctt tgcacgcagg aggtcagcgg ttcgatcccg ctattctcca ccatttttta	300
45	gagaaatggt gaaagattgc caagagacat tgtagtgag aatgaagaca caatgtctaa	360
47	tataagaaca atttaggttg tttttatatt agacttttta gtctaagttt atgttctaca	420
49	atttagaata cgacgctttg tgttggtgctg taggtttggg tctttaagat agctttgcta	480
51	tctggtgaaa gaacataaag atgttatatta atttattatt gtcaaagtca acaaacgca	540
53	aaaaaaacaa tttaacaactt gttagatggt ttacatttaa taaggagtg aaatgtgcat	600
55	tagaatacaa ataggtaagc tattaagagc gaatggtgga tgcctaggct gtaagaggcg	660
57	atgaaggacg tactagactg cgataagtta cggggagctg tcaagaagct ttgatccgta	720
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62 <210> SEQ ID NO: 2

63 <211> LENGTH: 527

64 <212> TYPE: DNA

65 <213> ORGANISM: Desulfitobacterium frappieri

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72	agatgaagtg aaacgggtca aagctggaga agtctgaaga gacttcgaaa tgccgaagag	180
74	gcaaagcagg ggaaatctgc ataagatgac cctgaaatcg agtcaaacct gttcaagcgc	240

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80 ataacatctg aaaaacctga atgtggcgga gacgtttggt caagctacta agggcgtagc 420
82 gtggatgcct aggcgctaag agtcgaagaa ggacgcggcg agcggcgaaa cgccacgggg 480
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97 acgtaatccg gtgggtgctca tgggtggaac gctgacagct acttctcgtc cgggtcccgt 180
99 ttctgtgcgg gatccgagga gttatatcgg tgcactgttg ggtcctgaga gaacacgcga 240
101 gtgttttgtc agcgacgatg atccgcgaaa caagaggaca tggttttcctt gcggtagggg 300
103 ttgtttgtgt ttgtttgaga actgcacagt ggacgcgagc atctttgttg taagtgttta 360
105 tgagcgtagc gtggatgcct tggcaccagg agccgatgaa ggacgtggga ggctgcgata 420
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112 <212> TYPE: DNA
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115 <400> SEQUENCE: 4
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134 <211> LENGTH: 952
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143 acataggtgg aaacatcatg atctggcatt ggcgggacac cgccgtcttc gtttctcttt 180
145 cttcgcggac aagcttgacg cccaggttgc ggtccttttg actgcgttcc ggtttcgggc 240
147 ctgtagctca ggtggttaga gcgcaccctt gataaggggtg aggtcggacg ttcgagtcgt 300
149 cccaggccca ccaccatcag acagttcttg cctgcgcctc atgtccgaag cttcgcgaac 360
151 tctcgcctgt ggcatcctgt gatggggcca tagctcagtt gggagagcgc gtgctttgca 420
153 agcatgaggt cgtcggttcg atcccgtctg gctccaccat tcttcttttc ttgaggaaga 480
155 tgatggcagg gtggtttgcg ctcggtcctt ttgagtgaag gctcttgggg tcttgagcgt 540
157 cttgtccgcg aatatctgtt tcgcatgttc catcatgccg gtctccggcg gaacatgcac 600
159 ggctgtatga catcgtgaat agggcattga tcgactgtac cgtggcaaca cggtcgggtc 660
161 gtggggaagg tggcgacacc tttcgatgcg atcattgggt gctgaccgca ccattgtcga 720
163 caatgcgaag ctggtctttt caaagaagac gtcgaagccg tccggccggg agcaatcctg 780

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167 attcgggtgga tgccttggcg ctaagaggcg aagaaggacg tgatacgctg cgataagctt      900
169 cggggagccg cgaatgggct ttgatccgga gatttccgaa tggggcaacc ca              952
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173 <211> LENGTH: 579
174 <212> TYPE: DNA
175 <213> ORGANISM: Mycobacterium L1
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182 atggcccttt cacctgtagt ggggtggggg ctgggtgcacg acaagcaaac gaccaggatg      180
184 gggaccttcc ttgtgggggt tgtctgggtc tgccaaacac actgttgggc tttagacaaa      240
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190 caaatgtggc tctcgaggct ttcgggtctg gggggtgtgt ttgtgtgctt ttgatgtgca      420
192 gtttcttttt tcgaattggg tttttgtgtt gtaagtgttt aagggcgcat ggtggatgcc      480
194 ttggcactgg gagccgatga aggacgtggg aggctgcgtt atgcctcggg gagctgtcaa      540
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200 <211> LENGTH: 523
201 <212> TYPE: DNA
202 <213> ORGANISM: Desulfomicrobium norvegicum
204 <400> SEQUENCE: 7
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209 gctcaggtgg ttagagcgca cgcctgataa gcgtgaggtc ggaagttcaa gtcttcccag      180
211 gcccaccatt tcttagtggt ggtgtagctc agctgggaga gcgcctgcct tgcacgcagg      240
213 aggtcatcag ttcgatcctg ttcacctcca ccattttcca actcgacaag aatttatgtt      300
215 gctagtcttt atcgtcagag tgtcttttga cactatggcg cccaagcata gcagcttggtg      360
217 atcattgaca gacgaatagg tgaagagaag agagttaaga tgtaagggc atacgggtgga      420
219 tgccttggcg tcaggaggcg atgaaggacg tggaaggctg cgataagcct cggggagccg      480
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224 <210> SEQ ID NO: 8
225 <211> LENGTH: 662
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234 gcaatgaagt gaaacgattc aaagtcggag aagtcttaag agacttctta taggaaactt      180
236 ggcttgtgtg aagcatgagc agaagccata gttgacttat ccacggagtg gaaaaatgcc      240
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246 aactgtaatt aggataacat ctaaaaccta gaagtggcgg caaaaaacgt ttggtcaagc      540
248 tactaagggc gtacgggtgga tgcctaggcg ctaagagtcg aagaaggacg cggcgagcgg      600
250 cgaaacgcca cggggagcag taagcatgcc ttgatccgtg gatatccgaa tggggcaacc      660
252 ca              662

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265 agatgaagtg aaacggttca aagctggaga agtctataga gacttcgaag tgccgaagag      180
267 gcaaagcagg ggaaatctgc ataagatgac cctgaagtcg agtcaaacct gttcaagcgc      240
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271 ctgggttagag cgcacgcctg ataagcgtga ggctcgggtgt tgcagtcacac ctaggcccac      360
273 cattattcaa agaggataga gacccgaacc tccaaacaat acttcacgcc agaacatacc      420
275 taacaggggt gagtattgag aggggagcgg ctccctctct aacgacatgg gggatatagct      480
277 cagctggggg agcacctgcc ttgcaagcag ggggtcagcg gttcgatccc gcttacctcc      540
279 accatcatat actggtttct ctaatgttct ttgaaaactg cacagagaag aaaaaactgt      600
281 aatttaggat aacatctgaa aaacctgaat gtggcggaga cggttggtca agctactaag      660
283 ggcgtacggt ggatgcctag gcgctaagag tcgaagaagg acgcggcgag cggcgaaacg      720
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288 <210> SEQ ID NO: 10
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290 <212> TYPE: DNA
291 <213> ORGANISM: Clostridium formicoaceticum
293 <400> SEQUENCE: 10
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298 aaaatcttag attttgtgtt agtcgcttag ttaaaaattc tgtaattcac gacaatagtt      180
300 ttaaaccaac aaaaaatgaa tggagaagatt tttaacatct atagtctttt agattgttct      240
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304 tattaagggc aaaggggtgga tgccttggca ctaggagccg aagaaggacg tggttaagctg      360
306 cgaaaagcca cggggagctg caagcaagta ttgatccgtg gatgtccgaa tggggaaacc      420
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312 <211> LENGTH: 699
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314 <213> ORGANISM: Desulfuromonas chloroethenica
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356 cacgaaaacg taaattatta ggatcaagaa gaaaagagca caggggtgaat gccttggcaa      300
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360 taaagcgtag atatccgaat ggggcaaccc a                                391
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365 <212> TYPE: DNA
366 <213> ORGANISM: Dehalobacter restrictus
368 <400> SEQUENCE: 13
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373 actggactga ctctcaagta aggtgagttt agcaatttat ttcttggtgt ttagttttga      180
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387 taacagcgaa atgccacggg gagtcgtaag caggcataga tccgtggatg tccgaatggg      600
389 gaaaccca                                608
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393 <211> LENGTH: 689
394 <212> TYPE: DNA
395 <213> ORGANISM: Desulfitobacterium sp. strain PCE1
397 <400> SEQUENCE: 14
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402 gcaatgaagt gaaacgattc aaagttggag aagtcttaag agacttctga aagccgaaga      180
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416 gtggcgggcaa aaaacgtttg gtcaagctac taaggcgcta cgggtggatgc ctaggcgcta      600
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420 atccgtggat atccgaatgg ggcaaccca                                689
423 <210> SEQ ID NO: 15
424 <211> LENGTH: 468
425 <212> TYPE: DNA
426 <213> ORGANISM: Desulfitobacterium frappieri TCE1
428 <400> SEQUENCE: 15

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/01/2006
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:116,117,118

VERIFICATION SUMMARY

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L:12 M:270 C: Current Application Number differs, Replaced Current Application Number